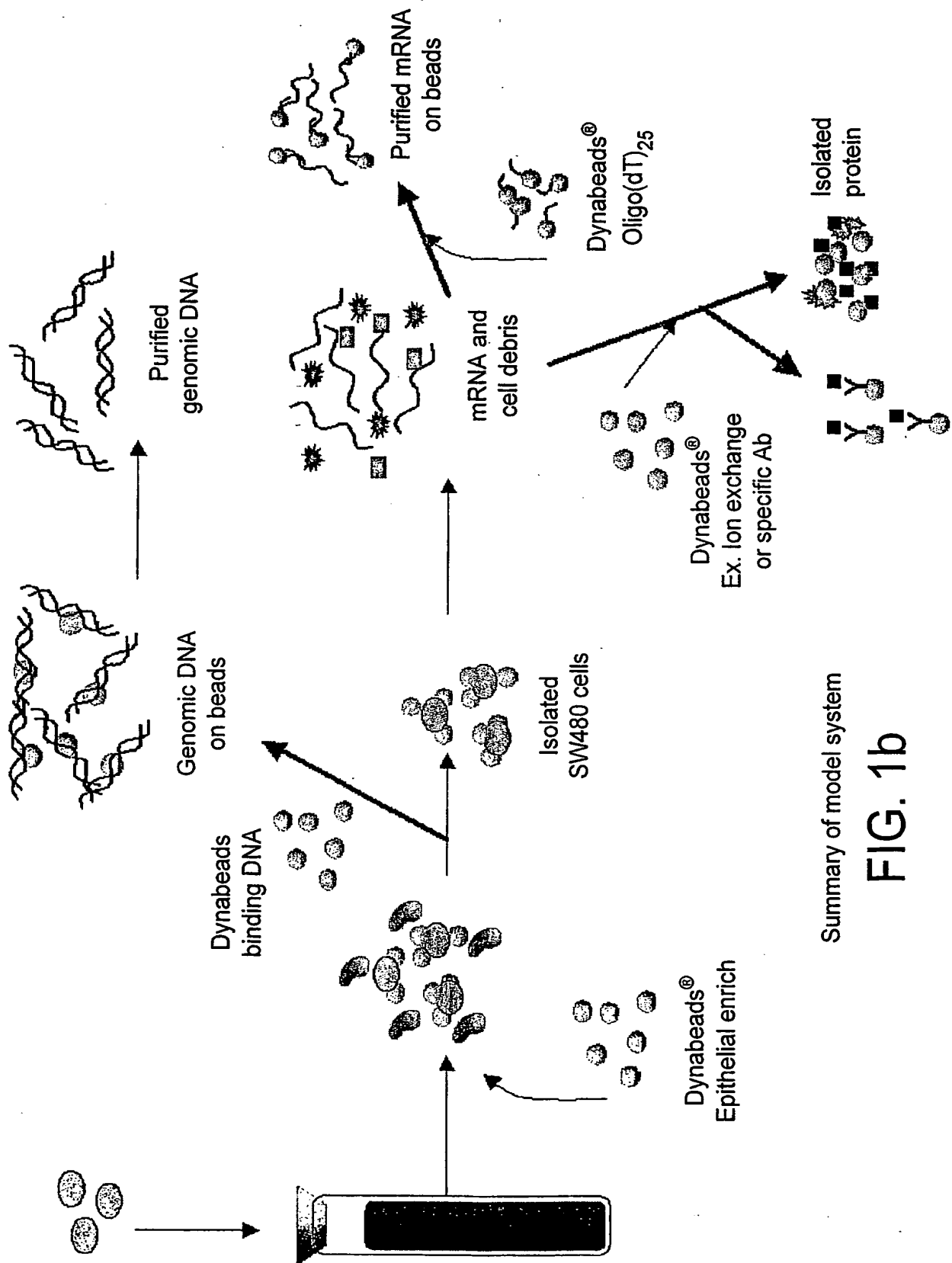


Summary of model system

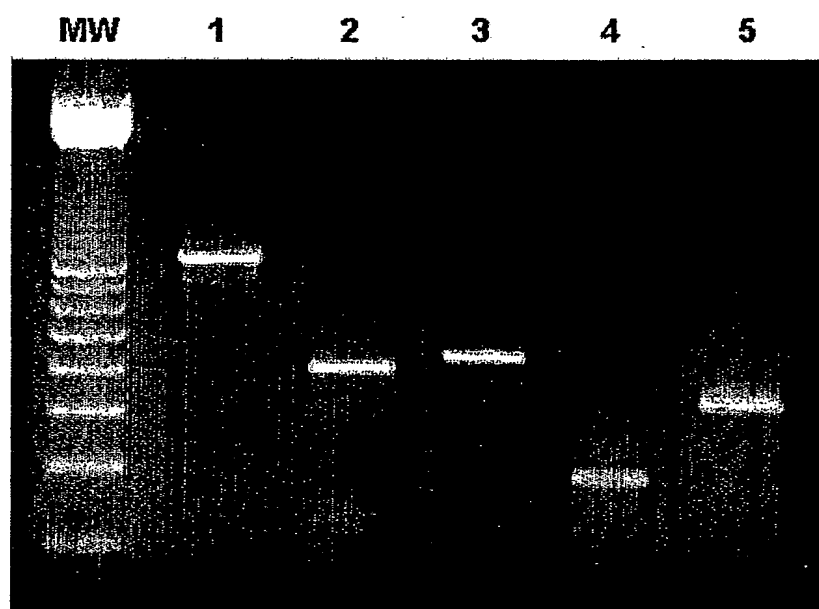
FIG. 1a

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Summary of model system

FIG. 1b



Re-use of Solid-phase
cDNA library in multiple
PCR experiments,
detecting different
transcripts

Cultured Carcinoma Cells

Lane	No. of cells	Amplicon	Round of PCR
1	10000	ESX	first (45 cycles)
2	10000	Mucin1	second (45 cycles)
3	10000	CK19	third (30 cycles)
4	10	CK19	first (nested)
5	10	ESX	second (nested)

FIG. 2

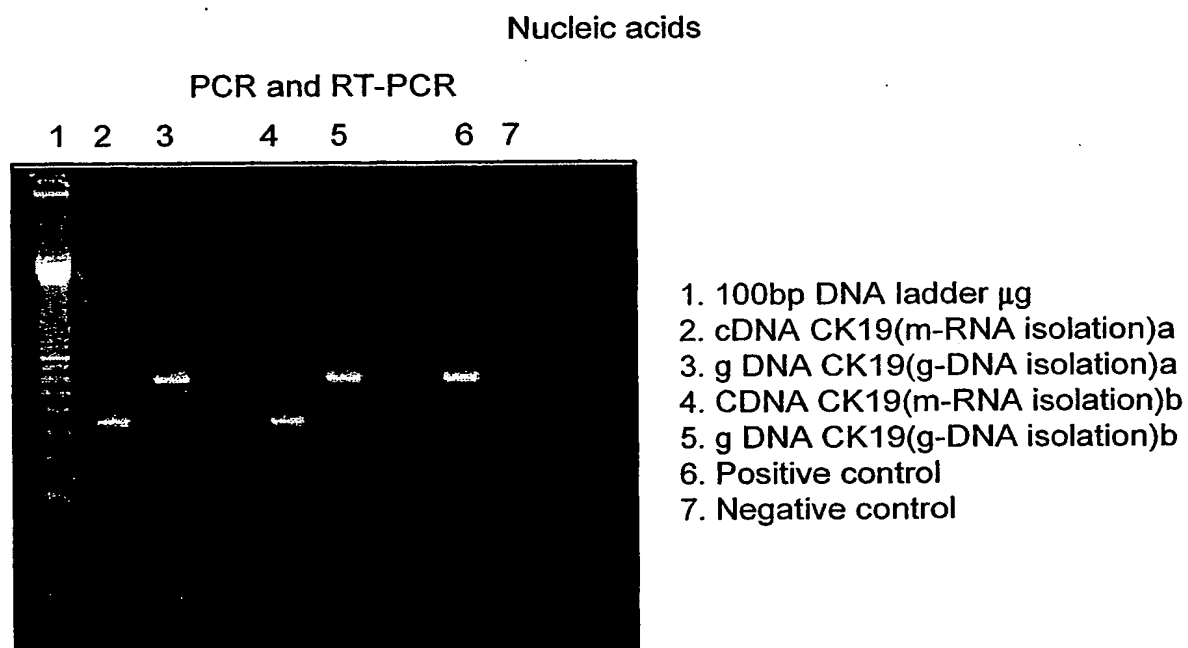


FIG. 3

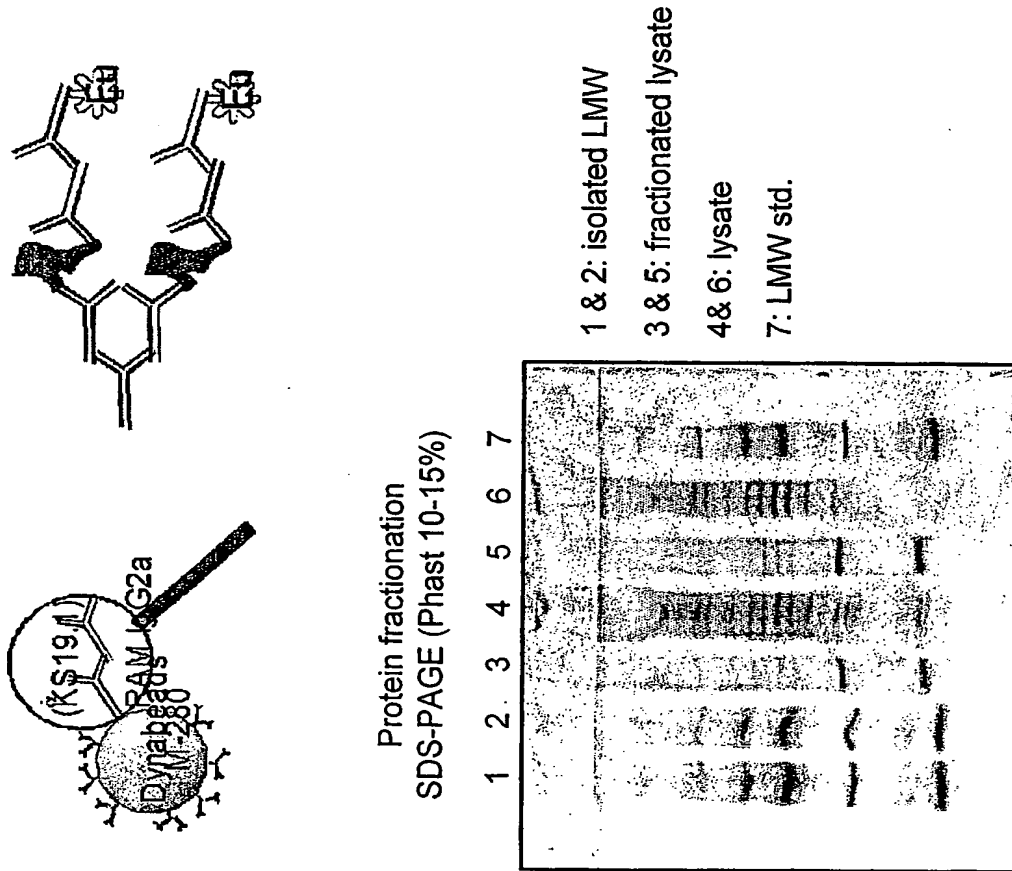
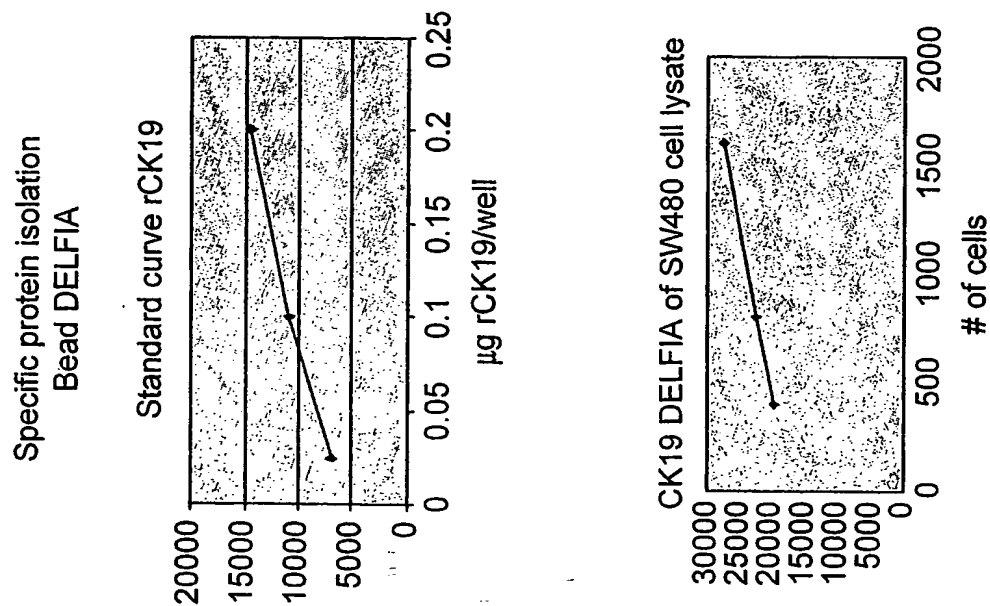
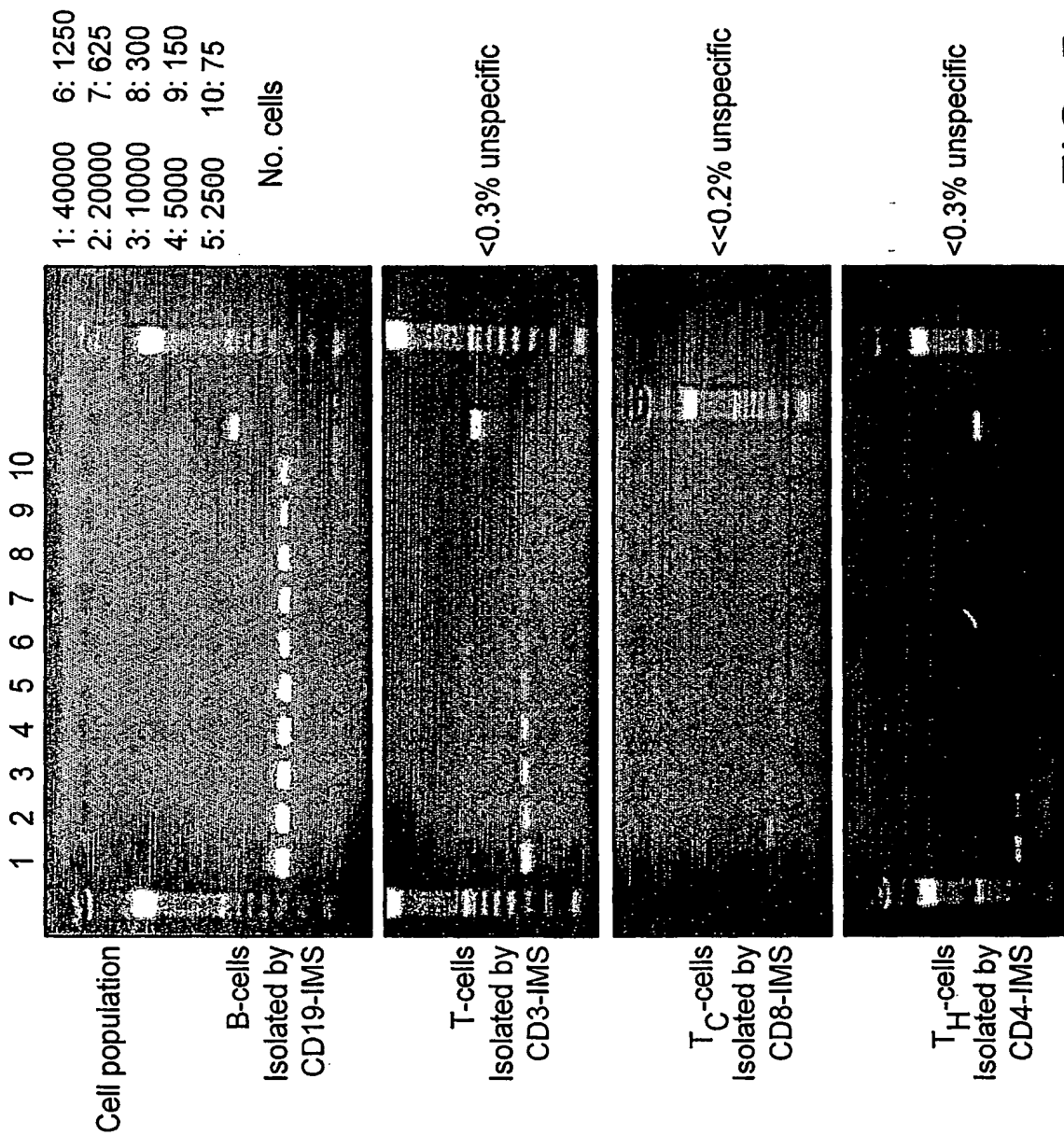


FIG. 4





Detection of B-cells by CD19 RT-PCR

FIG. 5a

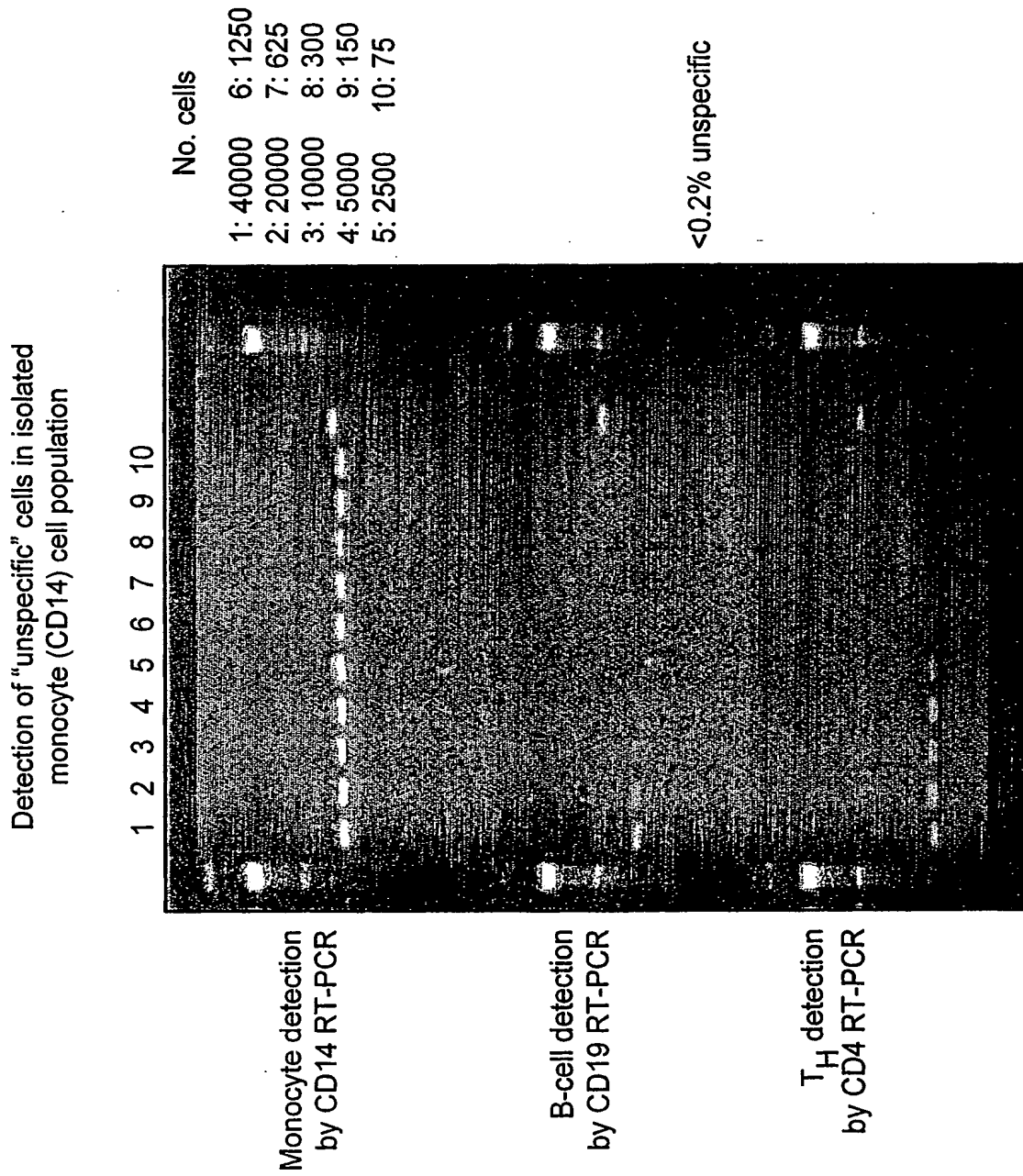
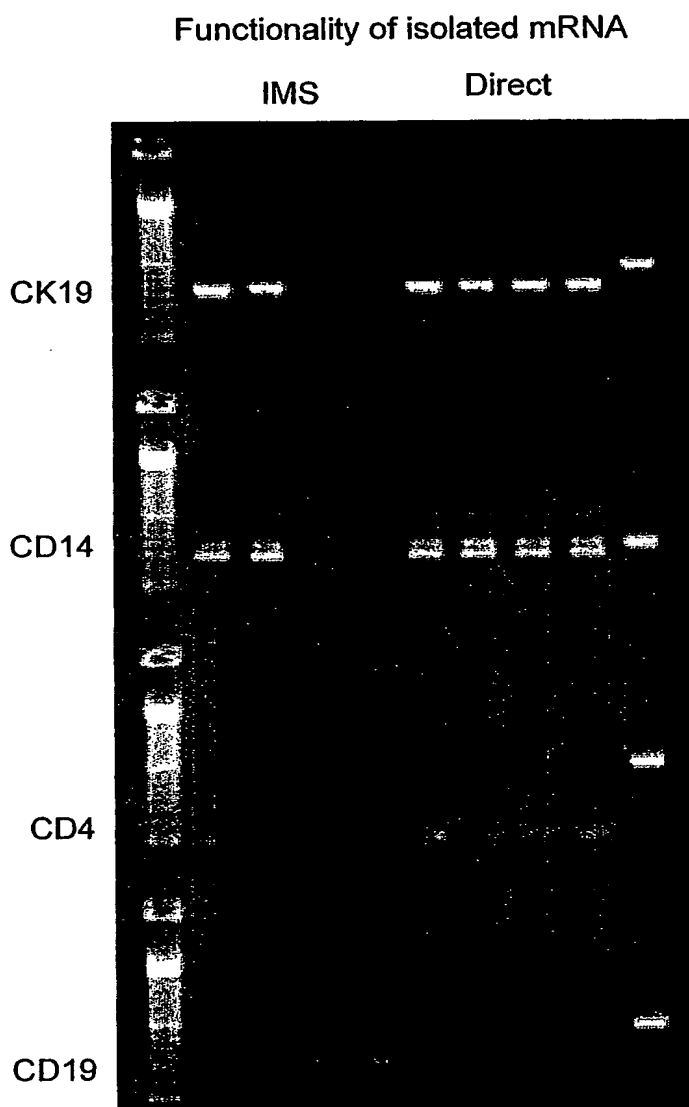
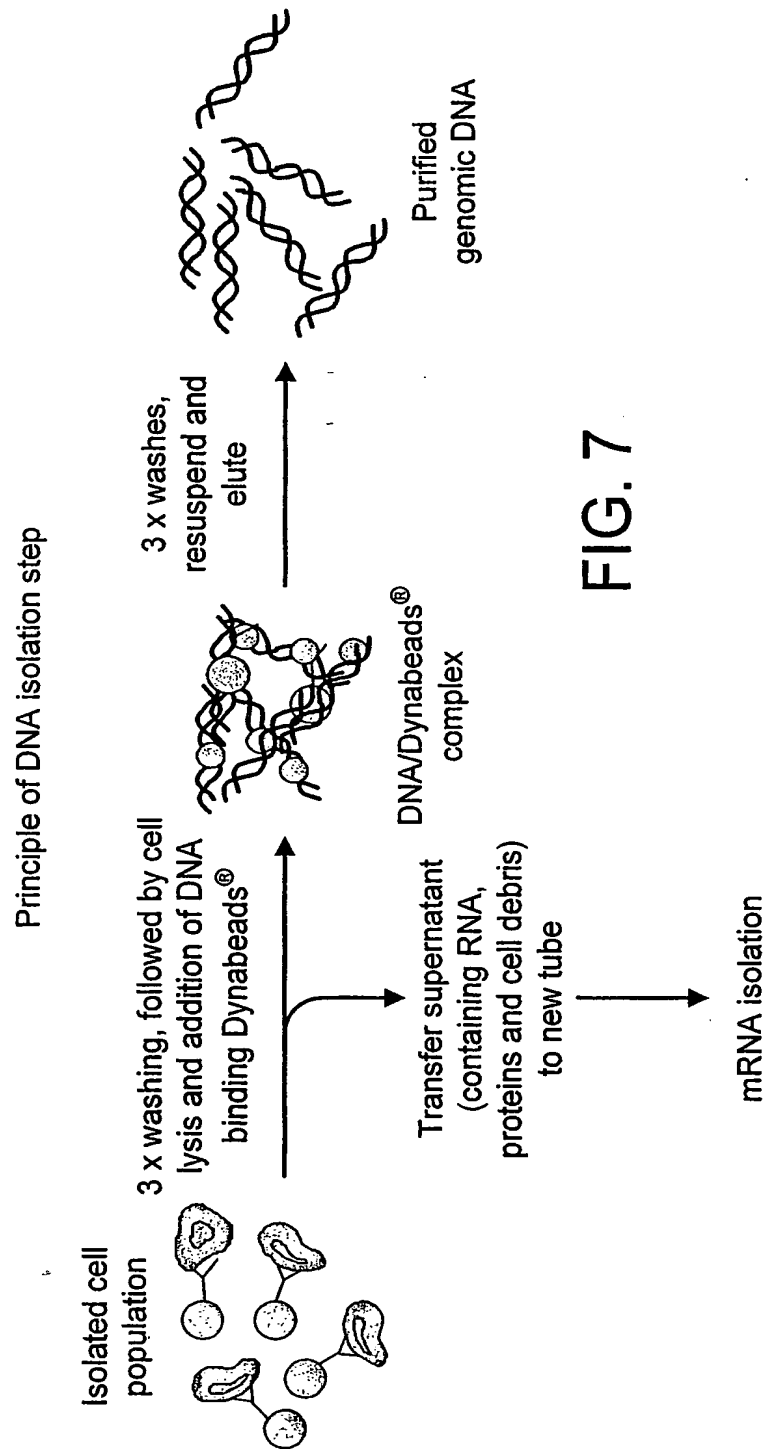


FIG. 5b

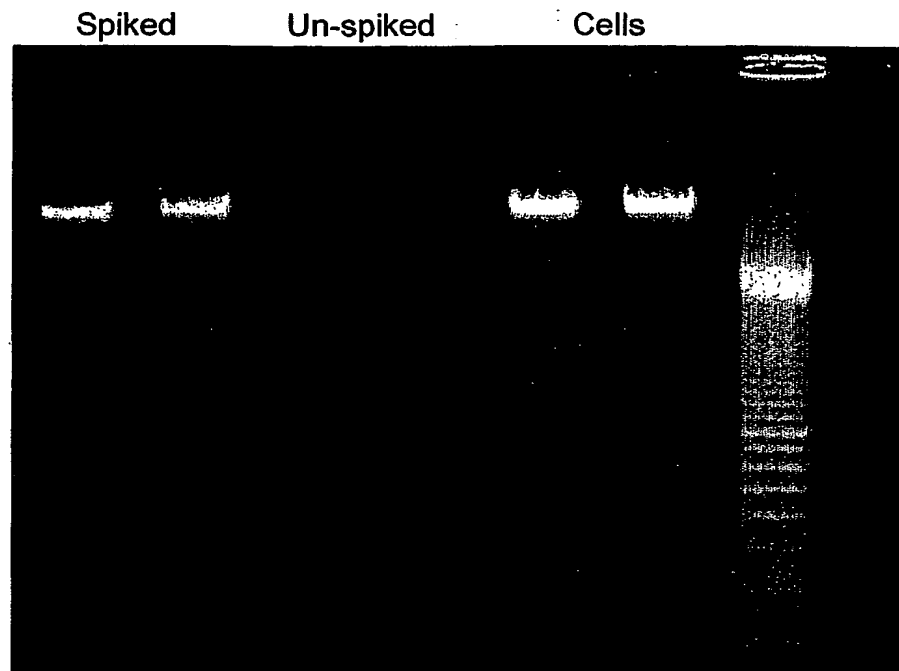


Template in RT-PCR:
mRNA representing
10000 carcinoma cells

FIG. 6



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Integrity of isolated DNA:
No degradation

FIG. 8

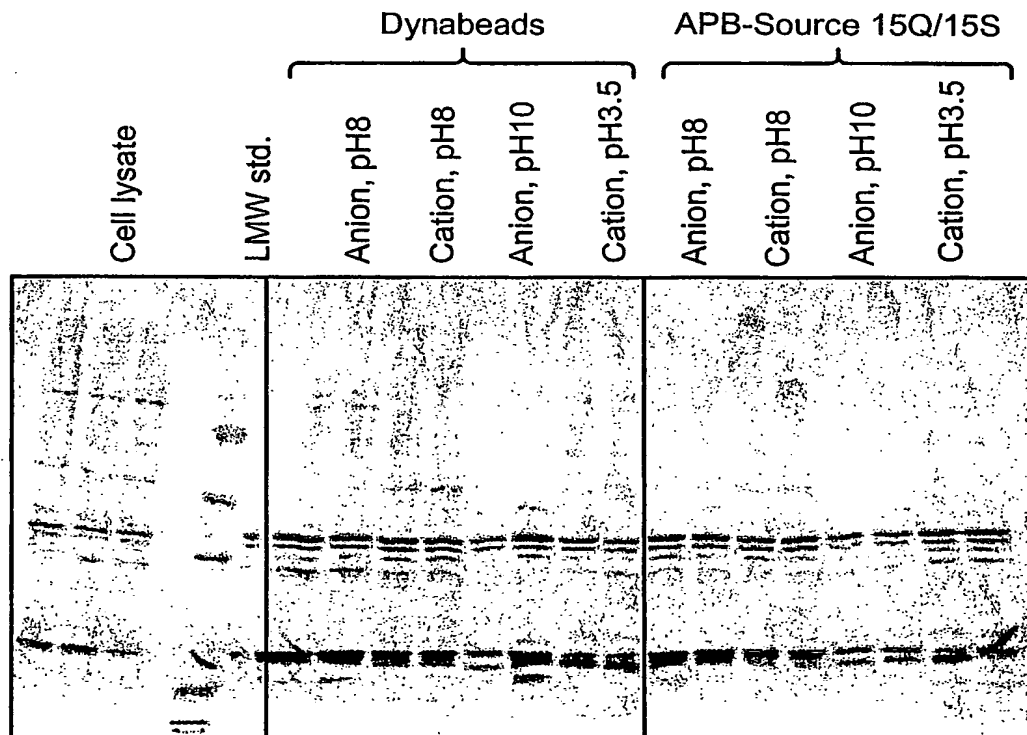


FIG. 9a

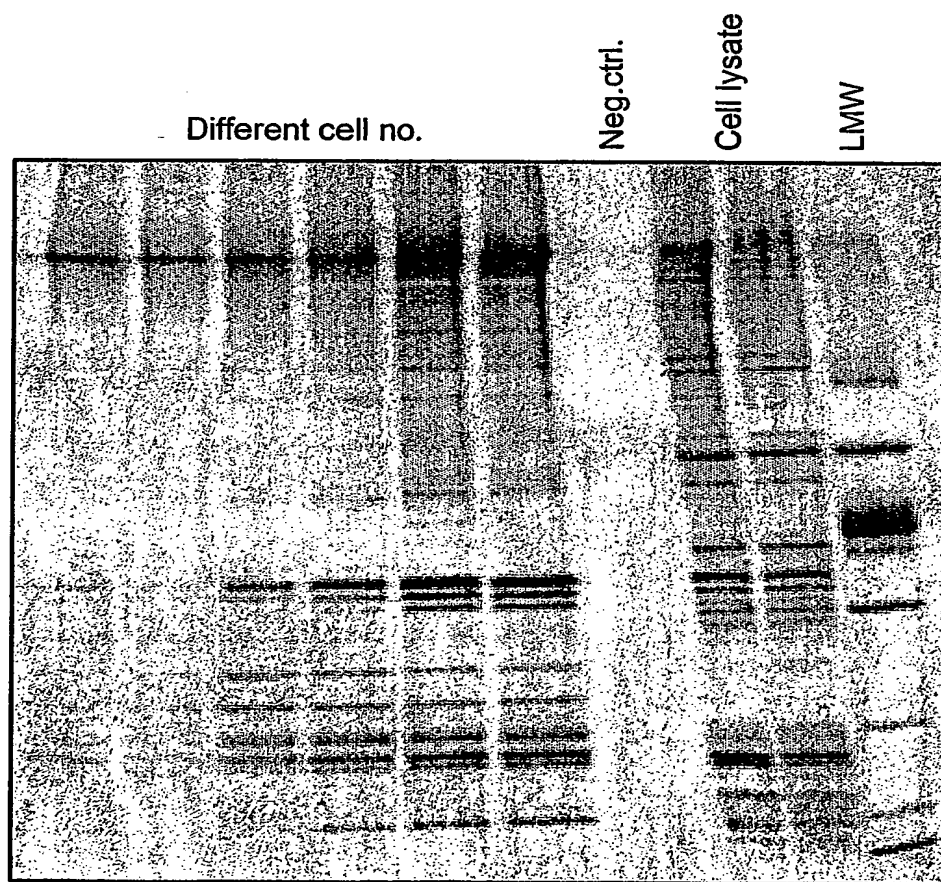
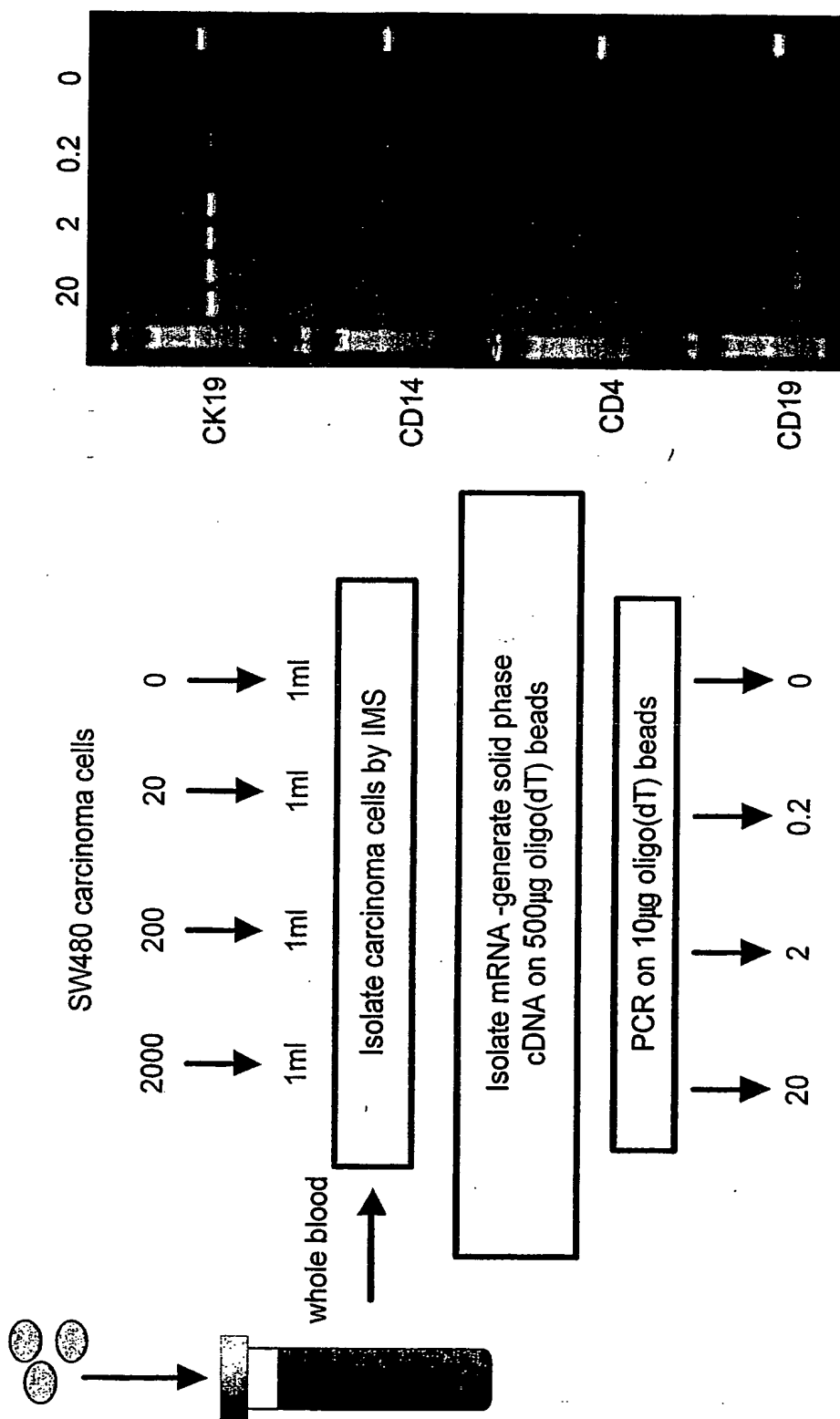


FIG. 9b



Representation of carcinoma cells in PCR

FIG. 10a

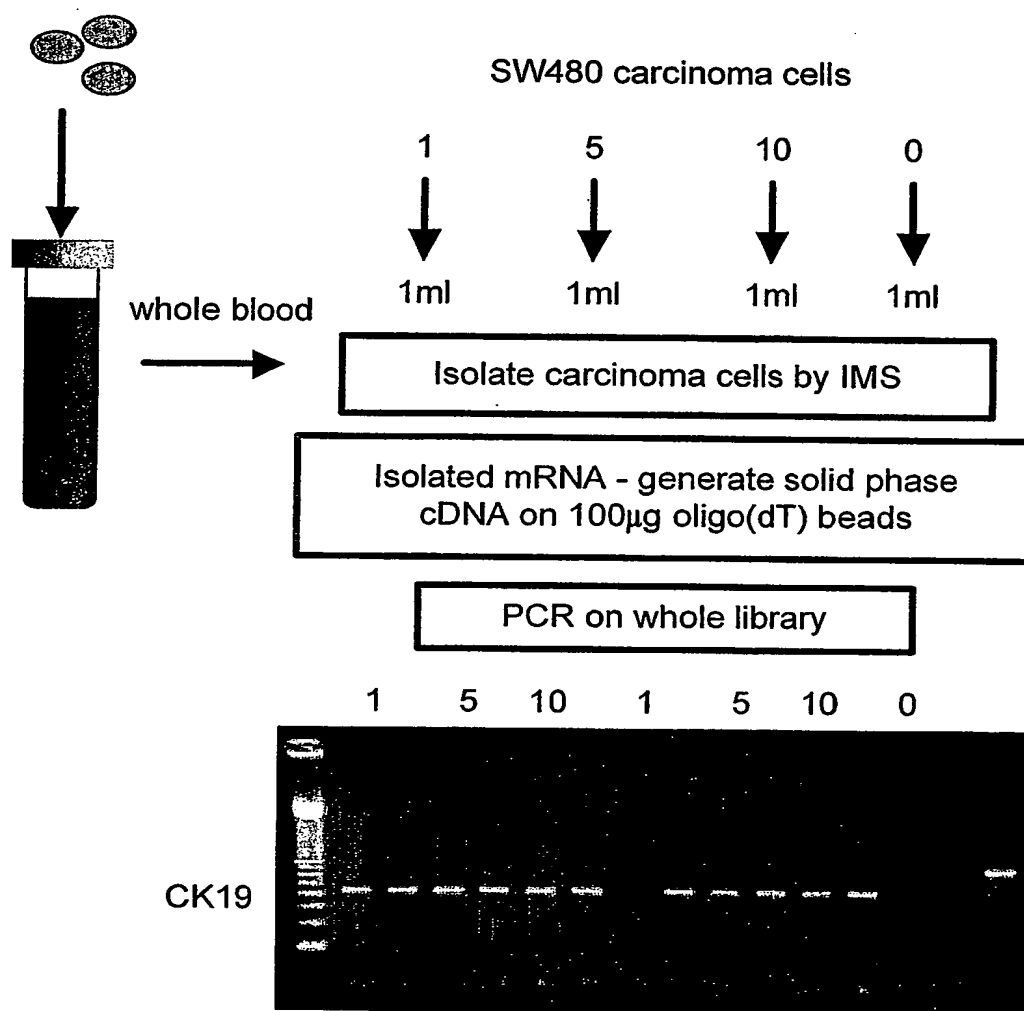


FIG. 10b

Flexible technology
Re-Use of Solid Phase cDNA

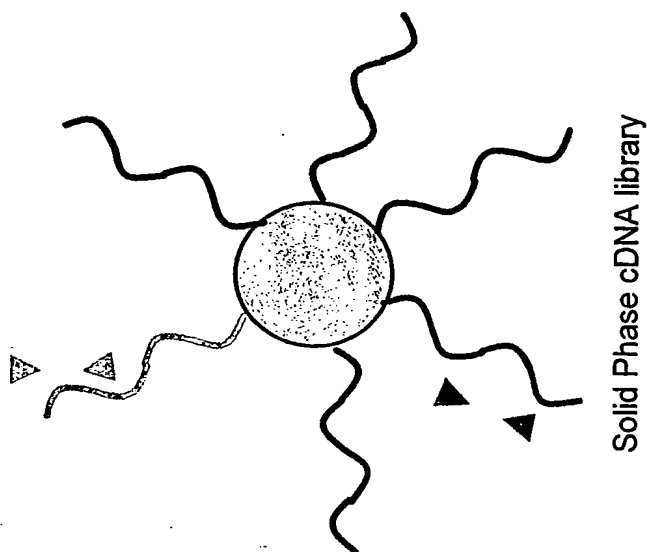
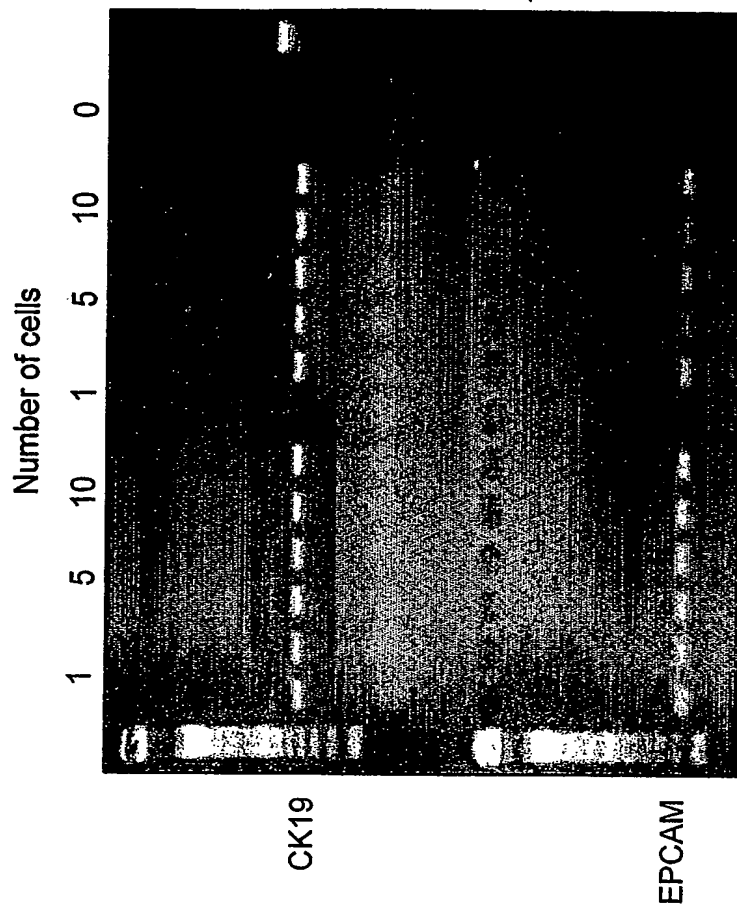
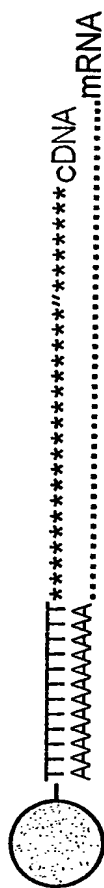


FIG. 11

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